=> s 14

SAMPLE SEARCH INITIATED 11:31:04 FILE 'REGISTRY'

SAMPLE SCREEN SEARCH COMPLETED - 0 TO ITERATE

100.0% PROCESSED 0 ITERATIONS 0 ANSWERS

SEARCH TIME: 00.00.01

FULL FILE PROJECTIONS: ONLINE **COMPLETE**

BATCH **COMPLETE**

PROJECTED ITERATIONS: 0 TO 0 PROJECTED ANSWERS: 0 TO 0

L5 0 SEA SSS SAM L4

=> s 14 ful

FULL SEARCH INITIATED 11:31:11 FILE 'REGISTRY'

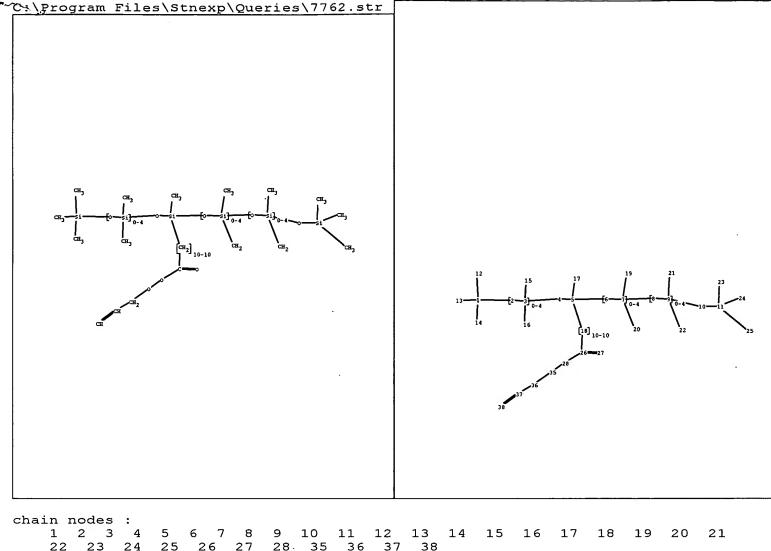
FULL SCREEN SEARCH COMPLETED - 1 TO ITERATE

100.0% PROCESSED 1 ITERATIONS 0 ANSWERS

SEARCH TIME: 00.00.01

L6 0 SEA SSS FUL L4

=>



22 23 24 25 26 27 28. 35 36 37 38 chain bonds : 2-3 3-4 3-15 3-16 4-5 5-6 5-17 5-18 6-7 7-8 1-2 1-12 1-13 1 - 147-19 7-20 8-9 9-10 9-21 9-22 10-11 11-23 11-24 11-25 18-26 26-27 26-28 28-35 35-36 36-37 37-38 exact/norm bonds : 26-27 26-28 exact bonds : 1-2 1-12 ·1-13 1-14 7-19 7-20 8-9 9-10 2-3 3-4 3-15 3-16 4-5 5-6 5-17 5-18 6-7 7-8 9-21 9-22 10-11 11-23 11-24 11-25 18-26 28-35 35-36 36-37 37-38

Match level : 1:CLASS 2:CLASS 3:CLASS 4:CLASS 5:CLASS 6:CLASS 7:CLASS 8:CLASS 9:CLASS 10:CLASS 11:CLASS 12:CLASS 13:CLASS 14:CLASS 15:CLASS 16:CLASS 17:CLASS 21:CLASS 22:CLASS 18:CLASS 19:CLASS 20:CLASS 28:CLASS 35:CLASS 23:CLASS 24:CLASS 25:CLASS 26:CLASS 27:CLASS 36:CLASS 37:CLASS 38:CLASS